



Entrez PubMed Nucleotide Protein Genome Structure PMC Taxonomy Books

Search Protein for Go Clear

Limits

Preview/Index

History

Clipboard

Details

Display

default

Show:

20

Send to

File

Get Subsequence

Fea

☐ 1: 1BKNB. Chain B, Crystal ...[gi:5107507]

BLink, Domains, Links

LOCUS 1BKN\_B 352 aa linear BCT 09-JUL-1998

DEFINITION Chain B, Crystal Structure Of An N-Terminal 40kd Fragment Of E.  
Coli Dna Mismatch Repair Protein MutL.

ACCESSION 1BKN\_B

VERSION 1BKN\_B GI:5107507

DBSOURCE pdb: molecule 1BKN, chain 66, release Jul 9, 1998;

deposition: Jul 9, 1998;

class: Dna Repair;

source: Mol\_id: 1; Organism\_scientific: Escherichia Coli; Strain:

K-12; Plasmid: Ptx418; Gene: MutL; Expression\_system: Escherichia

Coli; Expression\_system\_strain: Hms174 (Lambda De3);

Expression\_system\_vector\_type: Plasmid; Expression\_system\_plasmid:

Ptx418; Expression\_system\_gene: MutL;

Exp. method: X-Ray Diffraction.

## KEYWORDS

SOURCE Escherichia coli

ORGANISM Escherichia coli

Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;

Enterobacteriaceae; Escherichia.

REFERENCE 1 (residues 1 to 352)

AUTHORS Ban,C. and Yang,W.

TITLE Crystal structure and ATPase activity of MutL: implications for DNA  
repair and mutagenesis

JOURNAL Cell 95 (4), 541-552 (1998)

MEDLINE 99043508

PUBMED 9827806

REFERENCE 2 (residues 1 to 352)

AUTHORS Yang,W. and Ban,C.

TITLE Direct Submission

JOURNAL Submitted (09-JUL-1998)

COMMENT Revision History:

MAY 11 99 Initial Entry.

FEATURES Location/Qualifiers

source

1..352

/organism="Escherichia coli"

/db\_xref="taxon:562"

Region

1..222

/region\_name="Domain 3"

/note="NCBI Domains"

SecStr

26..40

/sec\_str\_type="helix"

/note="helix 7"

SecStr

41..51

/sec\_str\_type="sheet"

/note="strand 14"

SecStr

55..64

/sec\_str\_type="sheet"

/note="strand 15"

SecStr 97..106  
/sec\_str\_type="helix"  
/note="helix 8"  
SecStr 109..117  
/sec\_str\_type="sheet"  
/note="strand 16"  
SecStr 123..126  
/sec\_str\_type="sheet"  
/note="strand 17"  
SecStr 135..140  
/sec\_str\_type="sheet"  
/note="strand 18"  
SecStr 144..152  
/sec\_str\_type="sheet"  
/note="strand 19"  
SecStr 169..185  
/sec\_str\_type="helix"  
/note="helix 9"  
SecStr 187..195  
/sec\_str\_type="sheet"  
/note="strand 20"  
SecStr 196..205  
/sec\_str\_type="sheet"  
/note="strand 21"  
SecStr 211..219  
/sec\_str\_type="helix"  
/note="helix 10"  
Region 223..352  
/region\_name="Domain 4"  
/note="NCBI Domains"  
SecStr 226..235  
/sec\_str\_type="sheet"  
/note="strand 22"  
SecStr 236..245  
/sec\_str\_type="sheet"  
/note="strand 23"  
SecStr 256..262  
/sec\_str\_type="sheet"  
/note="strand 24"  
SecStr 269..282  
/sec\_str\_type="helix"  
/note="helix 11"  
SecStr 290..298  
/sec\_str\_type="sheet"  
/note="strand 25"  
SecStr 319..333  
/sec\_str\_type="helix"  
/note="helix 12"

## ORIGIN

1 gshmpiqvlp pqlanqiaag evverpasvv kelvenslda gatriddie rggaklirir  
61 dngcgikkde lalalarhat skiaslddle aiislgfrge alasissvsr ltitsrtaeq  
121 qeawqayaeg rdmnvtvkpa ahpvgttlev ldlfyntpar rkflrtekte fnhideiirr  
181 ialarfdvti nlshngkivr qyravpeggq kerrlgaicg taflegalai ewqhgdltr  
241 gwvadpnhtt palaeiqcy vngmmrdrl inhairqace dklgadqqa fvlyleidph  
301 qvdvnhpak hevrfhqsrl vhdfiyqgvl svlqqqletp lplddepqpa pr

//